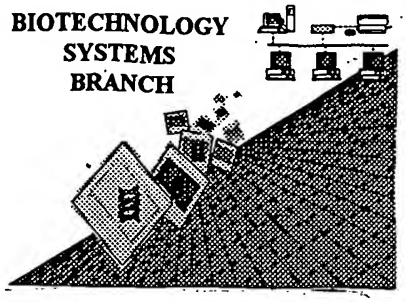
0400

BIOTECHNOLOGY SYSTEMS BRANCH



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/032,256
Source: OIPE
Date Processed by STIC: 1/16/2002

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO **REDUCE** ERRORED SEQUENCE LISTINGS, **PLEASE** USE THE **CHECKER VERSION 3.1 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
http://www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom, including:

1. EFS-Bio (<**http://www.uspto.gov/ebc/efs/downloads/documents.htm**>, **EFS Submission User Manual** - ePAVE)
2. U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. **Hand Carry directly to:**
   U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
   Or
   U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. **Federal Express, United Parcel Service, or other delivery service to:** U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: 10/032,256

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1	Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor **after** creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2	Invalid Line Length	The rules require that a line **not exceed** 72 characters in length. This includes white spaces.
3	Misaligned Amino Numbering	The numbering under each 5th amino acid is misaligned. Do **not** use tab codes between numbers; use **space characters**, instead.
4	Non-ASCII	The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. **Please ensure your subsequent submission is saved in ASCII text.**
5	Variable Length	Sequence(s)____contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6	PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**
7	Skipped Sequences (OLD RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for **each** skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.
8	Skipped Sequences (NEW RULES)	Sequence(s) _____ missing. If **intentional**, please insert the following lines for **each** skipped sequence. <210> sequence id number <400> sequence id number 000
9	Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of **n** or **Xaa**, and which residue **n** or **Xaa** represents.
10	Invalid <213> Response	Per 1.823 of Sequence Rules, the only **valid** <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence
11	Use of <220>	Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
12	PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
13	Misuse of n	n can only be used to represent a single **nucleotide** in a nucleic acid sequence. N is **not** used to represent any value not specifically a nucleotide.

OIPE

RAW SEQUENCE LISTING DATE: 01/16/2002
PATENT APPLICATION: US/10/032,256 TIME: 18:45:02

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Output Set: N:\CRF3\01162002\J032256.raw

Does Not Comply
Corrected Diskette Needed

P.5

3 <110> APPLICANT: CHODOSH, Lewis A
4 GARDNER, Heather P
6 <120> TITLE OF INVENTION: HORMONALLY UP-REGULATED, NEU-TUMOR-ASSOCIATED KINASE
8 <130> FILE REFERENCE: 22253-70421
C-1 10 **<140> CURRENT APPLICATION NUMBER: US/10/032,256**
C-1 11 **<141> CURRENT FILING DATE: 2001-12-21**
13 <150> PRIOR APPLICATION NUMBER: 60/257,073
14 <151> PRIOR FILING DATE: 2000-12-21
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22 <212> TYPE: DNA
23 <213> ORGANISM: Murinae gen. sp.
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28 ggggcgatgg aggcgcggag gacacgacca ggccggcggc ggcctgcgag ggaagtttcc 180
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31 gcgaggggct gcacgtgctg acgggagaaa aggtagctat caaggtcatc gataagaaaa 360
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49 aaaagcccaa agaacaagaa aaaagaggtg attttctcca ccgtccgttt tccaagaagt 1440
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PATENT APPLICATION: US/10/032,256 TIME: 18:45:02

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59 gttcaccgcc tcccgttccc agtaatggcc tcctgcagcc tctggggagc cccaactgtg 2040
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RAW SEQUENCE LISTING DATE: 01/16/2002
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<213> ORGANISM: Murinae gen. sp.

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            20                  25                  30
Glu Gly Ser Phe Leu Pro Ala Trp Val Ser Gly Val Ser Arg Glu Arg
        35                  40                  45
Leu Arg Asp Phe Gln His His Lys Arg Val Gly Asn Tyr Leu Ile Gly
    50                  55                  60
Ser Arg Lys Leu Gly Glu Gly Ser Phe Ala Lys Val Arg Glu Gly Leu
65                  70                  75                  80
His Val Leu Thr Gly Glu Lys Val Ala Ile Lys Val Ile Asp Lys Lys
                85                  90                  95
Arg Ala Lys Lys Asp Thr Tyr Val Thr Lys Asn Leu Arg Arg Glu Gly
            100                 105                 110
Gln Ile Gln Gln Met Ile Arg His Pro Asn Ile Thr Gln Leu Leu Asp
        115                 120                 125
Ile Leu Glu Thr Glu Asn Ser Tyr Tyr Leu Val Met Glu Leu Cys Pro
    130                 135                 140
Gly Gly Asn Leu Met His Lys Ile Tyr Glu Lys Lys Arg Leu Asp Glu
145                 150                 155                 160
Ala Glu Ala Arg Arg Tyr Ile Arg Gln Leu Ile Ser Ala Val Glu His
                165                 170                 175
Leu His Arg Ala Gly Val Val His Arg Asp Leu Lys Ile Glu Asn Leu
            180                 185                 190
Leu Leu Asp Glu Asp Asn Asn Ile Lys Leu Ile Asp Phe Gly Leu Ser
        195                 200                 205
Asn Cys Ala Gly Ile Leu Gly Tyr Ser Asp Pro Phe Ser Thr Gln Cys
    210                 215                 220
Gly Ser Pro Ala Tyr Ala Ala Pro Glu Leu Leu Ala Arg Lys Lys Tyr
225                 230                 235                 240
Gly Pro Lys Ile Asp Val Trp Ser Ile Gly Val Asn Met Tyr Ala Met
                245                 250                 255
Leu Thr Gly Thr Leu Pro Phe Thr Val Glu Pro Phe Ser Leu Arg Ala
            260                 265                 270
Leu Tyr Gln Lys Met Val Asp Lys Ala Met Asn Pro Leu Pro Thr Gln
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Leu Ser Thr Gly Ala Val Asn Phe Leu Arg Ser Leu Leu Glu Pro Asp
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PATENT APPLICATION: US/10/032,256 TIME: 18:45:02

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Output Set: N:\CRF3\01162002\J032256.raw

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176 305                 310                 315                 320
178 Asn Glu Asn Tyr Thr Gly Lys Val Pro Cys Asn Val Thr Tyr Pro Asn
179                 325                 330                 335
181 Arg Ile Ser Leu Glu Asp Leu Ser Pro Ser Val Val Leu His Met Thr
182             340                 345                 350
184 Glu Lys Leu Gly Tyr Lys Asn Ser Asp Val Ile Asn Thr Val Leu Ser
185         355                 360                 365
187 Asn Arg Ala Cys His Ile Leu Ala Ile Tyr Phe Leu Leu Asn Lys Lys
188     370                 375                 380
190 Leu Glu Arg Tyr Leu Ser Gly Lys Ser Asp Ile Gln Asp Ser Ile Cys
191 385                 390                 395                 400
193 Tyr Lys Thr Gln Leu Tyr Gln Ile Glu Lys Cys Arg Ala Thr Lys Glu
194                 405                 410                 415
196 Pro Tyr Glu Ala Ser Leu Asp Thr Trp Thr Arg Asp Phe Glu Phe His
197             420                 425                 430
199 Ala Val Gln Asp Lys Lys Pro Lys Glu Gln Glu Lys Arg Gly Asp Phe
200         435                 440                 445
202 Leu His Arg Pro Phe Ser Lys Lys Leu Asp Lys Asn Leu Pro Ser His
203     450                 455                 460
205 Lys Gln Pro Ser Pro Ser Leu Ile Thr Gln Leu Gln Ser Thr Lys Ala
206 465                 470                 475                 480
208 Leu Leu Lys Asp Arg Lys Ala Ser Lys Ser Gly Phe Pro Asp Lys Asp
209                 485                 490                 495
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212             500                 505                 510
214 Val Ala Ser Ser Ser Met Glu Phe Ile Pro Val Pro Pro Pro Arg Thr
215         515                 520                 525
217 Pro Arg Ile Val Lys Lys Leu Glu Pro His Gln Pro Gly Pro Gly Ser
218     530                 535                 540
220 Ala Ser Ile Leu Pro Lys Glu Glu Pro Leu Leu Leu Asp Met Val Arg
221 545                 550                 555                 560
223 Ser Phe Glu Ser Val Asp Arg Glu Asp His Ile Glu Leu Leu Ser Pro
224                 565                 570                 575
226 Ser His His Tyr Arg Ile Leu Ser Ser Pro Val Ser Leu Ala Arg Arg
227             580                 585                 590
229 Asn Ser Ser Glu Arg Thr Leu Ser Gln Gly Leu Leu Ser Gly Ser Thr
230         595                 600                 605
232 Ser Pro Leu Gln Thr Pro Leu His Ser Thr Leu Val Ser Phe Ala His
233     610                 615                 620
235 Glu Glu Lys Asn Ser Pro Pro Lys Glu Glu Gly Val Cys Ser Pro Pro
236 625                 630                 635                 640
238 Pro Val Pro Ser Asn Gly Leu Leu Gln Pro Leu Gly Ser Pro Asn Cys
239                 645                 650                 655
241 Val Lys Ser Arg Gly Arg Phe Pro Met Met Gly Ile Gly Gln Met Leu
242             660                 665                 670
244 Arg Lys Arg His Gln Ser Leu Gln Pro Ser Ser Glu Arg Ser Leu Asp
245         675                 680                 685
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RAW SEQUENCE LISTING DATE: 01/16/2002
PATENT APPLICATION: US/10/032,256 TIME: 18:45:02

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251 705                 710
254 <210> SEQ ID NO: 3
255 <211> LENGTH: 10
256 <212> TYPE: RNA
257 <213> ORGANISM: Unknown Organism
259 <220> FEATURE:
260 <223> OTHER INFORMATION: Description of Unknown Organism:Kozak consensus
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263 <400> SEQUENCE: 3
264 gccrccaugg                                                          10
267 <210> SEQ ID NO: 4
268 <211> LENGTH: 6
269 <212> TYPE: DNA
270 <213> ORGANISM: Unknown Organism
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315 <213> ORGANISM: Artificial Sequence
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see item 9 on Error Summary Sheet

VERIFICATION SUMMARY DATE: 01/16/2002
PATENT APPLICATION: US/10/032,256 TIME: 18:45:03

Input Set : A:\M2335hn1.app
Output Set: N:\CRF3\01162002\J032256.raw

L:10 M:270 C: Current Application Number differs, Replaced Application Number
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:309 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:7
L:309 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:7
L:309 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7